46

RAW SEQUENCE LISTING PATENT APPLICATION US/09/060,609

DATE: 04/23/98 TIME: 14:55:12

INPUT SET: S25254.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

	1			SEQ	UENCE LISTING	
	2					ENTERED
	3	(1) G	eneral Info	ormation:		
	4	, , ,	ADDI TOANS	0b B		•
	5 6	(1)	APPLICANT:	Ozenberger, Brad	A.	
	7			Jacobsen, J. S. Kajkowski, Eilee		
	8 ⁻			Rajkowski, Eliee	:11	
	9	/ii)	יידייניה סה ז	NVENTION. B-Amvlo	id Peptide-Binding	Proteins and
	10	(11)		cleotides Encoding		rioteins and
	11		rorynac	reotides Encoding	the bame	
	12	(111)	NUMBER OF	SEQUENCES: 2		
	13	(,				
	14	(iv)	CORRESPONI	ENCE ADDRESS:		
	15	` ,		SSEE: American Ho	me Products	
	16			T: One Campus Dri		
	17		(C) CITY:	Parsippany		
	18		(D) STATE	: NJ		
	19		(E) COUNT	RY: USA		
	20		(F) ZIP:	07054		
	21					
	22	(V)		READABLE FORM:		
	23		(A) MEDIU	JM TYPE: Floppy di	sk	
	24		• •	JTER: IBM PC compa		
	25		, <i>,</i>	ATING SYSTEM: PC-D		
	26		(D) SOFTW	NARE: PatentIn Rel	ease #1.0, Version	#1.30
	27					
	28	(Vi)		PPLICATION DATA:		
->	0/29			CATION NUMBER: US	i	
	30		(B) FILIN			
	31		(C) CLASS	SIFICATION:		
	32 33	/ *** i i i \	3 mmod New / 3	GENT INFORMATION:		
	34	(\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \		Walsh, Andrea C.		
	35			TRATION NUMBER: 3		
	36		• •	RENCE/DOCKET NUMBER		
	37		(C) KEIE	CENCE, DOCKET NOME		
	38	(ix)	TELECOMMUN	NICATION INFORMATI	ON:	
	39	(/		PHONE: 973-683-216		
	40			PAX: 973-683-4117		
	41		(-,			
	42					
	43	(2) INFO	RMATION FOR	R SEQ ID NO:1:		
	44	- •		*		
	45	(i)	SEQUENCE C	CHARACTERISTICS:		
		•				

(A) LENGTH: 810 base pairs

PAGE: 2 RAW SEQUENCE LISTING PATENT APPLICATION US/09/060,609

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47			(1	B) T	YPE:	nuc.	leic	aci	đ								
48	(C) STRANDEDNESS: single																
49			()	D) T(OPOL	OGY:	line	ear									
50																	
51		(ii) MO	LECU	LE T	YPE:	mRN	A									
52																	
53					_												
54		(ix) FE														
55			•	•		KEY:											
56 53			(1	8) T(JCAT.	ION:	1	80 /									
57 58																	
59		191	\ CF/	ALIENIA.	ית שי	ESCR	TDWT	.	GEV .	TT NI	٥.1.						
60		(V T) SE	SOFM	JE DI	BOCK.	TLIT	JN : .	SEQ .	או עד	0:1:						
61	ΔТС	САТ	ΔΨΨ	υψδ	ΔΔΔ	GGG	тст	aaa	ΔΔΤ	стс	ΔΨΨ	CCA	caa	GCT	CAC	aaa	48
62												Pro					10
63	1				5	1				10			5		15	1	
64	_									- •							
65	CAG	AAG	AAC	ACG	CGA	AGA	GAC	GGA	ACT	GGC	CTC	TAT	CCT	ATG	CGA	GGT	96
66	Gln	Lys	Asn	Thr	Arg	Arg	Asp	Gly	Thr	Gly	Leu	Tyr	Pro	Met	Arg	Gly	
67		-		20	_	_	_	_	25	_		_		30	_	_	
68																	
69	CCC	TTT	AAG	AAC	CTC	GCC	CTG	TTG	CCC	TTC	TCC	CTC	CCG	CTC	CTG	GGC	144
70	Pro	Phe	Lys	Asn	Leu	Ala	Leu	Leu	Pro	Phe	Ser	Leu	Pro	Leu	Leu	Gly	
71			35					40					45				
72																	
73												GTC					192
74	GΤÄ	_	GTĀ	Ser	СТÂ	Ser	_	GIu	Lys	Val	Ser	Val	Ser	Lys	Met	Ala	
75 76		50					55					60					
76 77	aaa	aaa	шаа	aaa	шаш	aam	aaa	mam	aam	aaa	asa	GCC	ama	3.00	000	303	240
78												Ala					240
79	65	AIG	пр	110	Ser	70	110	Der	ATO	FIO	75	ALG	Val	TIIL	лта	80	
80	0.5					, 0					, 5					00	
81	CTC	GTT	GGT	GTC	CTG	TGG	TTC	GTC	TCA	GTC	ACT	ACA	GGA	ccc	TGG	GGG	288
82												Thr					
83			•		85	•				90			•		95	-	
84																	
85	GCT	GTT	GCC	ACC	TCC	GCC	GGG	GGC	GAG	GAG	TCG	CTT	AAG	TGC	GAG	GAC	336
86	Ala	Val	Ala	Thr	Ser	Ala	Gly	Gly	Glu	Glu	Ser	Leu	Lys	Cys	Glu	Asp	
87				100					105					110			
88																	
89												AAA					384
90	Leu	Lys		Gly	Gln	Tyr	Ile	_	Lys	Asp	Pro	Lys		Asn	Asp	Ala	
91			115					120					125				
92				~~-	ae		m~=	. ~ .		m. ~	. ~ -	~~~	a	a==	ma	m~=	4.0.0
93												GCT					432
94	Tnr		GIU	Pro	vaı	asn	_	rnr	ASN	тyr	rnr	Ala	HIS	vaı	ser	cys	
95 96		130					135					140					
96 97	արա	CCA	CCA	מממ	אאמ	ηmη	אכייי	ጥረጥ	አአመ	CAT	שממ	AGT	aaa	חאא	GAA	አሮአ	480
9 <i>1</i> 98												Ser					400
99	145	110	WT.G	110	POII	150	1111	Cys	пyз	roh	155	Ser	G T Y	NOII	OIU	160	
,,	113										133					100	

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100																		
101	CAT	TTT	ACT	GGG	AAC	GAA	GTT	GGT	TTT	TTC	AAG	CCC	ATA	TCT	TGC	CGA		528
102	His	Phe	Thr	Gly	Asn	Glu	Val	Gly	Phe	Phe	Lys	Pro	Ile	Ser	Cys	Arg		
103					165					170					175			
104																		
105	AAT	GTA	AAT	GGC	TAT	TCC	TAC	AAA	GTG	GCA	GTC	GCA	TTG	TCT	CTT	TTT		576
106	Asn	Val	Asn	Gly	Tyr	Ser	Tyr	Lys	Val	Ala	Val	Ala	Leu	Ser	Leu	Phe		
107				180	_		_	_	185					190				
108																		
109	CTT	GGA	TGG	TTG	GGA	GCA	GAT	CGA	TTT	TAC	CTT	GGA	TAC	CCT	GCT	TTG		624
110	Leu	Gly	Trp	Leu	Gly	Ala	Asp	Arg	Phe	Tyr	Leu	Gly	Tyr	Pro	Ala	Leu		
111		_	195		_		_	200		_		_	205					
112																		
113	GGT	TTG	TTA	AAG	TTT	TGC	ACT	GTA	GGG	TTT	TGT	GGA	ATT	GGG	AGC	CTA		672
114	Gly	Leu	Leu	Lys	Phe	Cys	Thr	Val	Gly	Phe	Cys	Gly	Ile	Gly	Ser	Leu		
115		210		_		_	215		_		_	220		_				
116																		
117	ATT	GAT	TTC	ATT	CTT	ATT	TCA	ATG	CAG	ATT	GTT	GGA	CCT	TCA	GAT	GGA		720
118	Ile	Asp	Phe	Ile	Leu	Ile	Ser	Met	Gln	Ile	Val	Gly	Pro	Ser	Asp	Gly		
119	225					230					235					240		
120																		
121	AGT	AGT	TAC	ATT	ATA	GAT	TAC	TAT	GGA	ACC	AGA	CTT	ACA	AGA	CTG	AGT		768
122	Ser	Ser	Tyr	Ile	Ile	Asp	Tyr	Tyr	Gly	Thr	Arg	Leu	Thr	Arg	Leu	Ser		
123					245					250	_			_	255			
124																		
125	ATT	ACT	AAT	GAA	ACA	TTT	AGA	AAA	ACG	CAA	TTA	TAT	CCA	TAA				810
126	Ile	Thr	Asn	Glu	Thr	Phe	Arg	Lys	Thr	Gln	Leu	Tyr	Pro					
127				260					265		•							
128																		
129																		
130	(2)	INFO	RMAT	NOI	FOR	SEQ	ID N	NO:2:	:									
131																		
132		((i) S	SEQUE	ENCE	CHAI	RACTI	ERIST	rics:	}								
133				(A)	LEN	IGTH:	269	ami	ino a	acids	3							
134				(B)	TYE	E: a	amino	aci	id									
135				(D)	TOE	OLOC	3Y:]	linea	ar									

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met His Ile Leu Lys Gly Ser Pro Asn Val Ile Pro Arg Ala His Gly

Gln Lys Asn Thr Arg Arg Asp Gly Thr Gly Leu Tyr Pro Met Arg Gly

Pro Phe Lys Asn Leu Ala Leu Leu Pro Phe Ser Leu Pro Leu Leu Gly

Gly Gly Gly Ser Gly Ser Gly Glu Lys Val Ser Val Ser Lys Met Ala

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153 154 155	Ala 65	Ala	Trp	Pro	Ser	Gly 70	Pro	Ser	Ala	Pro	Glu 75	Ala	Val	Thr	Ala	Arg 80
156 157 158	Leu	Val	Gly	Val	Leu 85	Trp	Phe	Val	Ser	Val 90	Thr	Thr	Gly	Pro	Trp 95	Gly
159 160 161	Ala	Val	Ala	Thr 100	Ser	Ala	Gly	Gly	Glu 105	Glu	Ser	Leu	Lys	Cys 110	Glu	Asp
162 163 164	Leu	Lys	Val 115	Gly	Gln	Tyr	Ile	Cys 120	Lys	Asp	Pro	Lys	Ile 125	Asn	Asp	Ala
165 166 167	Thr	Gln 130	Glu	Pro	Val	Asn	Cys 135	Thr	Asn	Tyr	Thr	Ala 140	His	Val	Ser	Cys
168 169 170	Phe 145	Pro	Ala	Pro	Asn	Ile 150	Thr	Cys	Lys	Asp	Ser 155	Ser	Gly	Asn	Glu	Thr 160
171 172 173	His	Phe	Thr	Gly	Asn 165	Glu	Val	Gly	Phe	Phe 170	Lys	Pro	Ile	Ser	Cys 175	Arg
174 175 176	Asn	Val	Asn	Gly 180	Tyr	Ser	Tyr	Lys	Val 185	Ala	Val	Ala	Leu	Ser 190	Leu	Phe
177 178 179	Leu	Gly	Trp 195	Leu	Gly	Ala	Asp	Arg 200	Phe	Tyr	Leu	Gly	Tyr 205	Pro	Ala	Leu
180 181 182	Gly	Leu 210	Leu	Lys	Phe	Cys	Thr 215	Val	Gly	Phe	Cys	Gly 220	Ile	Gly	Ser	Leu
183 184 185	Ile 225	Asp	Phe	Ile	Leu	Ile 230	Ser	Met	Gln	Ile	Val 235	Gly	Pro	Ser	Asp	Gly 240
186 187 188	Ser	Ser	Tyr	Ile	Ile 245	Asp	Tyr	Tyr	Gly	Thr 250	Arg	Leu	Thr	Arg	Leu 255	Ser
189 190 191	Ile	Thr	Asn	Glu 260	Thr	Phe	Arg	Lys	Thr 265	Gln	Leu	Tyr	Pro			

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PAGE: 1

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/060,609

DATE: 04/23/98 TIME: 14:55:18

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Original Text

29

Wrong application Serial Number

(A) APPLICATION NUMBER: US